RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _____ 10/719,587 A_____

Source: _____ IFW 16_____

Date Processed by STIC: _____ 08/17/2006_____

ENTERED

RAW SEQUENCE LISTING   DATE: 08/17/2006
PATENT APPLICATION: **US/10/719,587A**   TIME: 11:06:15

Input Set : **A:\55862-CIP(46342)-Amended.txt**
Output Set: **N:\CRF4\08172006\J719587A.raw**

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      3 <110> APPLICANT: Takeda Chemical Industries, Ltd.
      5 <120> TITLE OF INVENTION: Novel G protein-coupled receptor protein, its DNA and ligand
thereof
      7 <130> FILE REFERENCE: 2568US0P-CIP
C-->  9 <140> CURRENT APPLICATION NUMBER: US/10/719,587A
C-->  9 <141> CURRENT FILING DATE: 2003-11-21
      9 <150> PRIOR APPLICATION NUMBER: US 09/831,758
     10 <151> PRIOR FILING DATE: 2001-05-11
     12 <150> PRIOR APPLICATION NUMBER: PCT/JP99/06283
     13 <151> PRIOR FILING DATE: 1999-11-11
     15 <150> PRIOR APPLICATION NUMBER: JP 10-323759
     16 <151> PRIOR FILING DATE: 1998-11-13
     18 <150> PRIOR APPLICATION NUMBER: JP 11-060030
     19 <151> PRIOR FILING DATE: 1999-03-08
     21 <150> PRIOR APPLICATION NUMBER: JP 11-106812
     22 <151> PRIOR FILING DATE: 1999-04-14
     24 <150> PRIOR APPLICATION NUMBER: JP 11-166672
     25 <151> PRIOR FILING DATE: 1999-06-14
     27 <150> PRIOR APPLICATION NUMBER: JP 11-221640
     28 <151> PRIOR FILING DATE: 1999-08-04
     30 <150> PRIOR APPLICATION NUMBER: JP 11-259818
     31 <151> PRIOR FILING DATE: 1999-09-14
     33 <160> NUMBER OF SEQ ID NOS: 66
     35 <210> SEQ ID NO: 1
     36 <211> LENGTH: 180
     37 <212> TYPE: PRT
     38 <213> ORGANISM: Human
     40 <400> SEQUENCE: 1
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     42 1               5                   10                  15
     43 Ser Ser Leu Leu Thr Ser Asn Ile Phe Cys Ala Asp Glu Leu Val Met
     44             20                  25                  30
     45 Ser Asn Leu His Ser Lys Glu Asn Tyr Asp Lys Tyr Ser Glu Pro Arg
     46         35                  40                  45
     47 Gly Tyr Pro Lys Gly Glu Arg Ser Leu Asn Phe Glu Glu Leu Lys Asp
     48     50                  55                  60
     49 Trp Gly Pro Lys Asn Val Ile Lys Met Ser Thr Pro Ala Val Asn Lys
     50 65                  70                  75                  80
     51 Met Pro His Ser Phe Ala Asn Leu Pro Leu Arg Phe Gly Arg Asn Val
     52                 85                  90                  95
     53 Gln Glu Glu Arg Ser Ala Gly Ala Thr Ala Asn Leu Pro Leu Arg Ser
     54             100                 105                 110
     55 Gly Arg Asn Met Glu Val Ser Leu Val Arg Arg Val Pro Asn Leu Pro
     56             115                 120                 125
```

```
57 Gln Arg Phe Gly Arg Thr Thr Thr Ala Lys Ser Val Cys Arg Met Leu
58         130             135             140
59 Ser Asp Leu Cys Gln Gly Ser Met His Ser Pro Cys Ala Asn Asp Leu
60 145                 150             155                 160
61 Phe Tyr Ser Met Thr Cys Gln His Gln Glu Ile Gln Asn Pro Asp Gln
62                 165             170             175
63 Lys Gln Ser Arg
64               180
66 <210> SEQ ID NO: 2
67 <211> LENGTH: 540
68 <212> TYPE: DNA
69 <213> ORGANISM: Human
71 <400> SEQUENCE: 2
72 atggaaatta tttcatcaaa actattcatt ttattgactt tagccacttc aagcttgtta   60
73 acatcaaaca tttttgtgc agatgaatta gtgatgtcca atcttcacag caaagaaat  120
74 tatgacaaat attctgagcc tagaggatac ccaaaagggg aaagaagcct caattttgag  180
75 gaattaaaag attggggacc aaaaaatgtt attaagatga gtacacctgc agtcaataaa  240
76 atgccacact ccttcgccaa cttgccattg agatttggga ggaacgttca agaagaaga  300
77 agtgctggag caacagccaa cctgcctctg agatctgga agaaatatgga ggtgagcctc  360
78 gtgagacgtg ttcctaacct gccccaaagg tttgggagaa caacaacagc caaaagtgtc  420
79 tgcaggatgc tgagtgattt gtgtcaagga tccatgcatt caccatgtgc caatgactta  480
80 ttttactcca tgacctgcca gcaccaagaa atccagaatc ccgatcaaaa acagtcaagg  540
82 <210> SEQ ID NO: 3
83 <211> LENGTH: 27
84 <212> TYPE: DNA
85 <213> ORGANISM: Artificial Sequence
87 <220> FEATURE:
88 <223> OTHER INFORMATION: primer
90 <400> SEQUENCE: 3
91 gggctgcaca tagagactta attttag                                       27
93 <210> SEQ ID NO: 4
94 <211> LENGTH: 27
95 <212> TYPE: DNA
96 <213> ORGANISM: Artificial Sequence
98 <220> FEATURE:
99 <223> OTHER INFORMATION: primer
101 <400> SEQUENCE: 4
102 ctagaccacc tctatataac tgcccat                                       27
104 <210> SEQ ID NO: 5
105 <211> LENGTH: 30
106 <212> TYPE: DNA
107 <213> ORGANISM: Artificial Sequence
109 <220> FEATURE:
110 <223> OTHER INFORMATION: primer
112 <400> SEQUENCE: 5
113 gcacatagag acttaatttt agatttagac                                    30
115 <210> SEQ ID NO: 6
116 <211> LENGTH: 27
117 <212> TYPE: DNA
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118 <213> ORGANISM: Artificial Sequence
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121 <223> OTHER INFORMATION: primer
123 <400> SEQUENCE: 6
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126 <210> SEQ ID NO: 7
127 <211> LENGTH: 27
128 <212> TYPE: DNA
129 <213> ORGANISM: Artificial Sequence
131 <220> FEATURE:
132 <223> OTHER INFORMATION: primer
134 <400> SEQUENCE: 7
135 cagctttagg gacaggctcc aggtttc                                    27
137 <210> SEQ ID NO: 8
138 <211> LENGTH: 196
139 <212> TYPE: PRT
140 <213> ORGANISM: Human
142 <400> SEQUENCE: 8
143 Met Glu Ile Ile Ser Ser Lys Leu Phe Ile Leu Leu Thr Leu Ala Thr
144  1               5                   10                  15
145 Ser Ser Leu Leu Thr Ser Asn Ile Phe Cys Ala Asp Glu Leu Val Met
146             20                  25                  30
147 Ser Asn Leu His Ser Lys Glu Asn Tyr Asp Lys Tyr Ser Glu Pro Arg
148             35                  40                  45
149 Gly Tyr Pro Lys Gly Glu Arg Ser Leu Asn Phe Glu Glu Leu Lys Asp
150         50                  55                  60
151 Trp Gly Pro Lys Asn Val Ile Lys Met Ser Thr Pro Ala Val Asn Lys
152 65                  70                  75                  80
153 Met Pro His Ser Phe Ala Asn Leu Pro Leu Arg Phe Gly Arg Asn Val
154             85                  90                  95
155 Gln Glu Glu Arg Ser Ala Gly Ala Thr Ala Asn Leu Pro Leu Arg Ser
156             100                 105                 110
157 Gly Arg Asn Met Glu Val Ser Leu Val Arg Arg Val Pro Asn Leu Pro
158             115                 120                 125
159 Gln Arg Phe Gly Arg Thr Thr Thr Ala Lys Ser Val Cys Arg Met Leu
160         130                 135                 140
161 Ser Asp Leu Cys Gln Gly Ser Met His Ser Pro Cys Ala Asn Asp Leu
162 145                 150                 155                 160
163 Phe Tyr Ser Met Thr Cys Gln His Gln Glu Ile Gln Asn Pro Asp Gln
164             165                 170                 175
165 Lys Gln Ser Arg Arg Leu Leu Phe Lys Lys Ile Asp Asp Ala Glu Leu
166             180                 185                 190
167 Lys Gln Glu Lys
168             195
170 <210> SEQ ID NO: 9
171 <211> LENGTH: 588
172 <212> TYPE: DNA
173 <213> ORGANISM: Human
175 <400> SEQUENCE: 9
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176 atggaaatta tttcatcaaa actattcatt ttattgactt tagccacttc aagcttgtta   60
177 acatcaaaca ttttttgtgc agatgaatta gtgatgtcca atcttcacag caaagaaaat  120
178 tatgacaaat attctgagcc tagaggatac ccaaaagggg aaagaagcct caattttgag  180
179 gaattaaaag attggggacc aaaaaatgtt attaagatga gtacacctgc agtcaataaa  240
180 atgccacact ccttcgccaa cttgccattg agatttggga ggaacgttca agaagaaaga  300
181 agtgctggag caacagccaa cctgcctctg agatctggaa gaaatatgga ggtgagcctc  360
182 gtgagacgtg ttcctaacct gccccaaagg tttgggagaa caacaacagc caaaagtgtc  420
183 tgcaggatgc tgagtgattt gtgtcaagga tccatgcatt caccatgtgc caatgactta  480
184 ttttactcca tgacctgcca gcaccaagaa atccagaatc ccgatcaaaa acagtcaagg  540
185 agactgctat tcaagaaaat agatgatgca gaattgaaac aagaaaaa               588
187 <210> SEQ ID NO: 10
188 <211> LENGTH: 27
189 <212> TYPE: DNA
190 <213> ORGANISM: Artificial Sequence
192 <220> FEATURE:
193 <223> OTHER INFORMATION: primer
195 <400> SEQUENCE: 10
196 gcctagagga gatctaggct gggagga                                27
198 <210> SEQ ID NO: 11
199 <211> LENGTH: 27
200 <212> TYPE: DNA
201 <213> ORGANISM: Artificial Sequence
203 <220> FEATURE:
204 <223> OTHER INFORMATION: primer
206 <400> SEQUENCE: 11
207 gggaggaaca tggaagaaga aaggagc                                27
209 <210> SEQ ID NO: 12
210 <211> LENGTH: 27
211 <212> TYPE: DNA
212 <213> ORGANISM: Artificial Sequence
214 <220> FEATURE:
215 <223> OTHER INFORMATION: primer
217 <400> SEQUENCE: 12
218 gatggtgaat gcatggactg ctggagc                                27
220 <210> SEQ ID NO: 13
221 <211> LENGTH: 27
222 <212> TYPE: DNA
223 <213> ORGANISM: Artificial Sequence
225 <220> FEATURE:
226 <223> OTHER INFORMATION: primer
228 <400> SEQUENCE: 13
229 ttcctcccaa atctcagtgg caggttg                                27
231 <210> SEQ ID NO: 14
232 <211> LENGTH: 196
233 <212> TYPE: PRT
234 <213> ORGANISM: Bovine
236 <400> SEQUENCE: 14
237 Met Glu Ile Ile Ser Leu Lys Arg Phe Ile Leu Leu Met Leu Ala Thr
238  1               5                   10                  15
```

```
239 Ser Ser Leu Leu Thr Ser Asn Ile Phe Cys Thr Asp Glu Ser Arg Met
240         20                  25              30
241 Pro Asn Leu Tyr Ser Lys Lys Asn Tyr Asp Lys Tyr Ser Glu Pro Arg
242         35                  40              45
243 Gly Asp Leu Gly Trp Glu Lys Glu Arg Ser Leu Thr Phe Glu Glu Val
244     50                  55              60
245 Lys Asp Trp Ala Pro Lys Ile Lys Met Asn Lys Pro Val Val Asn Lys
246 65                  70                  75                  80
247 Met Pro Pro Ser Ala Ala Asn Leu Pro Leu Arg Phe Gly Arg Asn Met
248                 85                  90                  95
249 Glu Glu Glu Arg Ser Thr Arg Ala Met Ala His Leu Pro Leu Arg Leu
250             100                 105             110
251 Gly Lys Asn Arg Glu Asp Ser Leu Ser Arg Trp Val Pro Asn Leu Pro
252         115                 120             125
253 Gln Arg Phe Gly Arg Thr Thr Thr Ala Lys Ser Ile Thr Lys Thr Leu
254     130                 135             140
255 Ser Asn Leu Leu Gln Gln Ser Met His Ser Pro Ser Thr Asn Gly Leu
256 145                 150                 155                 160
257 Leu Tyr Ser Met Ala Cys Gln Pro Gln Glu Ile Gln Asn Pro Gly Gln
258                 165             170                 175
259 Lys Asn Leu Arg Arg Arg Gly Phe Gln Lys Ile Asp Asp Ala Glu Leu
260             180             185                 190
261 Lys Gln Glu Lys
262         195
264 <210> SEQ ID NO: 15
265 <211> LENGTH: 588
266 <212> TYPE: DNA
267 <213> ORGANISM: Bovine
269 <400> SEQUENCE: 15
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271 acatcaaaca tcttctgcac agacgaatca aggatgccca atctttacag caaaaagaat  120
272 tatgacaaat attccgagcc tagaggagat ctaggctggg agaagaaag aagtcttact  180
273 tttgaagaag taaaagattg ggctccaaaa attaagatga ataaacctgt agtcaacaaa  240
274 atgccacctt ctgcagccaa cctgccactg agatttggga ggaacatgga agaagaaagg  300
275 agcactaggg cgatggccca cctgcctctg agactcggaa aaaatagaga ggacagcctc  360
276 tccagatggg tcccaaatct gccccagagg tttggaagaa caacaacagc caaaagcatt  420
277 accaagaccc tgagtaattt gctccagcag tccatgcatt caccatctac caatgggcta  480
278 ctctactcca tggcctgcca gccccaagaa atccagaatc ctggtcaaaa gaacctaagg  540
279 agacggggat ccagaaaat agatgatgca gaattgaaac aagaaaaa             588
281 <210> SEQ ID NO: 16
282 <211> LENGTH: 27
283 <212> TYPE: DNA
284 <213> ORGANISM: Artificial Sequence
286 <220> FEATURE:
287 <223> OTHER INFORMATION: primer
289 <400> SEQUENCE: 16
290 ccctggggct tcttctgtct tctatgt                                     27
292 <210> SEQ ID NO: 17
293 <211> LENGTH: 26
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 08/17/2006
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:20; N Pos. 3,9
Seq#:21; N Pos. 3,9,12
Seq#:22; N Pos. 3,6,9
Seq#:23; N Pos. 3,6,9,12
Seq#:24; N Pos. 3,6,9
Seq#:25; N Pos. 3,6,9,12
Seq#:63; Xaa Pos. 10

Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:64,65,66

VERIFICATION SUMMARY   DATE: 08/17/2006
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L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:373 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:399 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:425 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:456 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:482 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:513 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:1145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63 after pos.:0